

SEQUENCE LISTING

<110> Bradfield, Christopher A.
Dolwick, Kristin M.
Carver, Lucy A.

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Detecting Agonists to the Ah Receptor

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gcg	ata	gct	act	cca	ctt	cag	cca	cca	tcc	ata	ctt	gaa	atc	cgg	acc	1228
Ala	Ile	Ala	Thr	Pro	Leu	Gln	Pro	Pro	Ser	Ile	Leu	Glu	Ile	Arg	Thr	
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aaa	aat	ttt	atc	ttt	aga	acc	aaa	cac	aaa	cta	gac	ttc	aca	cct	att	1276
Lys	Asn	Phe	Ile	Phe	Arg	Thr	Lys	His	Lys	Leu	Asp	Phe	Thr	Pro	Ile	
		285					290					295				
ggt	tgt	gat	gcc	aaa	gga	aga	att	gtt	tta	gga	tat	act	gaa	gca	gag	1324
Gly	Cys	Asp	Ala	Lys	Gly	Arg	Ile	Val	Leu	Gly	Tyr	Thr	Glu	Ala	Glu	
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ctg	tgc	acg	aga	ggc	tca	ggt	tat	cag	ttt	att	cat	gca	gct	gat	atg	1372
Leu	Cys	Thr	Arg	Gly	Ser	Gly	Tyr	Gln	Phe	Ile	His	Ala	Ala	Asp	Met	

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Gly	Met	Ile	Val	Phe	Arg	Leu	Leu	Thr	Lys	Asn	Asn	Arg	Trp	Thr	Trp																
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gtc	cag	tct	aat	gca	cgc	ctg	ctt	tat	aaa	aat	gga	aga	cca	gat	tat	1516															
Val	Gln	Ser	Asn	Ala	Arg	Leu	Leu	Tyr	Lys	Asn	Gly	Arg	Pro	Asp	Tyr																
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Ile	Ile	Val	Thr	Gln	Arg	Pro	Leu	Thr	Asp	Glu	Glu	Gly	Thr	Glu	His																
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Leu	Arg	Lys	Arg	Asn	Thr	Lys	Leu	Pro	Phe	Met	Phe	Thr	Thr	Gly	Glu																
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gct	gtg	ttg	tat	gag	gca	acc	aac	cct	ttt	cct	gcc	ata	atg	gat	ccc	1660															
Ala	Val	Leu	Tyr	Glu	Ala	Thr	Asn	Pro	Phe	Pro	Ala	Ile	Met	Asp	Pro																
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aca	tcc	act	cta	agc	aag	gac	tct	ctc	aat	cct	agt	tcc	ctc	ctg	gct	1756															
Thr	Ser	Thr	Leu	Ser	Lys	Asp	Ser	Leu	Asn	Pro	Ser	Ser	Leu	Leu	Ala																
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gcc	atg	atg	caa	caa	gat	gag	tct	att	tat	ctc	tat	cct	gct	tca	agt	1804															
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Phe	Ala	Gly	Gly	His	Pro	Gly	Leu	Phe	Gln	Asp	Ser	Lys	Asn	Ser	Asp																
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Leu	Tyr	Ser	Ile	Met	Lys	Asn	Leu	Gly	Ile	Asp	Phe	Glu	Asp	Ile	Arg																
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gat Asp	tct Ser	tta Leu	agt Ser 590	aag Lys	tct Ser	ccc Pro	ttc Phe	ata Ile 595	cct Pro	tca Ser	gat Asp	tat Tyr	caa Gln 600	cag Gln	caa Gln	2188
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caa Gln	gag Glu	ttc Phe 685	ccc Pro	tac Tyr	aaa Lys	tct Ser	gaa Glu 690	atg Met	gat Asp	tct Ser	atg Met	cct Pro 695	tat Tyr	aca Thr	cag Gln	2476
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aca Thr 715	gag Glu	ctg Leu	gac Asp	tac Tyr	cct Pro	atg Met	ggg Gly	agt Ser	ttt Phe	gaa Glu 725	cca Pro	tcc Ser	cca Pro	tac Tyr	ccc Pro 730	2572
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aca Thr	tgt Cys	tat Tyr 765	gct Ala	ggg Gly	gcc Ala	gtg Val	tcg Ser	atg Met	tat Tyr	cag Gln	tgc Cys	cag Gln	cca Pro	gaa Glu	cct Pro	2716
cag Gln	cac His	acc Thr	cac His	gtg Val	ggt Gly	cag Gln	atg Met	cag Gln	tac Tyr	aat Asn	cca Pro	gta Val	ctg Leu	cca Pro	ggc Gly	2764

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Gln Gln Ala Phe Leu Asn Lys Phe Gln Asn Gly Val Leu Asn Glu Thr			
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tat cca gct gaa tta aat aac ata aat aac act cag act acc aca cat			2860
Tyr Pro Ala Glu Leu Asn Asn Ile Asn Asn Thr Gln Thr Thr Thr His			
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ctt cag cca ctt cat cat ccg tca gaa gcc aga cct ttt cct gat ttg			2908
Leu Gln Pro Leu His His Pro Ser Glu Ala Arg Pro Phe Pro Asp Leu			
	830	835	840
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Thr Ser Ser Gly Phe Leu			
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Ser Asn Pro Ser Lys Arg His Arg Asp Arg Leu Asn Thr Glu Leu Asp
          35          40          45
Arg Leu Ala Ser Leu Leu Pro Phe Pro Gln Asp Val Ile Asn Lys Leu
          50          55          60
Asp Lys Leu Ser Val Leu Arg Leu Ser Val Ser Tyr Leu Arg Ala Lys
          65          70          75          80
Ser Phe Phe Asp Val Ala Leu Lys Ser Ser Pro Thr Glu Arg Asn Gly
          85          90          95
Gly Gln Asp Asn Cys Arg Ala Ala Asn Phe Arg Glu Gly Leu Asn Leu

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Gln	Glu	Gly	Glu	Phe	Leu	Leu	Gln	Ala	Leu	Asn	Gly	Phe	Val	Leu	Val
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Tyr	Leu	Gly	Phe	Gln	Gln	Ser	Asp	Val	Ile	His	Gln	Ser	Val	Tyr	Glu
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Leu	Ile	His	Thr	Glu	Asp	Arg	Ala	Glu	Phe	Gln	Arg	Gln	Leu	His	Trp
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Ala	Leu	Asn	Pro	Ser	Gln	Cys	Thr	Glu	Ser	Gly	Gln	Gly	Ile	Glu	Glu
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Arg	Cys	Leu	Leu	Asp	Asn	Ser	Ser	Gly	Phe	Leu	Ala	Met	Asn	Phe	Gln
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Gly	Lys	Leu	Lys	Tyr	Leu	His	Gly	Gln	Lys	Lys	Lys	Gly	Lys	Asp	Gly
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305					310					315					320
Gly	Tyr	Gln	Phe	Ile	His	Ala	Ala	Asp	Met	Leu	Tyr	Cys	Ala	Glu	Ser
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His	Ile	Arg	Met	Ile	Lys	Thr	Gly	Glu	Ser	Gly	Met	Ile	Val	Phe	Arg
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Pro	Leu	Thr	Asp	Glu	Glu	Gly	Thr	Glu	His	Leu	Arg	Lys	Arg	Asn	Thr
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Lys	Leu	Pro	Phe	Met	Phe	Thr	Thr	Gly	Glu	Ala	Val	Leu	Tyr	Glu	Ala
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Thr	Asn	Pro	Phe	Pro	Ala	Ile	Met	Asp	Pro	Leu	Pro	Leu	Arg	Thr	Lys
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<400> 11
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<210> 13
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<212> DNA
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<210> 14
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<400> 15
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<212> DNA
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<220>
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Glu Gly Ile Lys Ser Asn Pro Ser Lys Arg His Arg Asp Arg Leu Asn

20 25 30
 Thr Glu Leu Asp Arg Leu Ala Ser Leu Leu Pro Phe Pro Gln Asp Val
 35 40 45
 Ile Asn Lys Leu Asp Lys Leu Ser Val Leu Arg Leu Ser Val Ser Tyr
 50 55 60
 Leu Arg Ala Lys Ser Phe Phe Asp
 65 70

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 1 5 10 15
 Glu Gly Ile Lys Ser Asn Pro Ser Lys Arg His Arg Asp Arg Leu Asn
 20 25 30
 Thr Glu Leu Asp Arg Leu Ala Ser Leu Leu Pro Phe Pro Gln Asp Val
 35 40 45
 Ile Asn Lys Leu Asp Lys Leu Ser Val Leu Arg Leu Ser Val Thr Tyr
 50 55 60
 Leu Arg Ala Lys Ser Phe Phe Asp
 65 70

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 20 25 30
 Val Cys Tyr Asn Pro Asp Gln Ile Pro Pro Glu Asn Ser Pro Leu Met
 35 40 45
 Glu Arg Cys Phe Ile Cys Arg Leu Arg Cys Leu Leu Asp Asn Ser Ser
 50 55 60
 Gly Phe Leu Ala Met Asn Phe Gln Gly Lys Leu Lys Tyr Leu His Gly
 65 70 75 80

Gln Lys Lys Lys Gly Lys Asp Gly Ser Ile Leu Pro Pro Gln Leu Ala
 85 90 95
 Leu Phe Ala Ile Ala Thr Pro Leu Gln Pro Pro Ser Ile Leu Glu Ile
 100 105 110
 Arg Thr Lys Asn Phe Ile Phe Arg Thr Lys His Lys Leu Asp Phe Thr
 115 120 125
 Pro Ile Gly Cys Asp Ala Lys Gly Arg Ile Val Leu Gly Tyr Thr Glu
 130 135 140
 Ala Glu Leu Cys Thr Arg Gly Ser Gly Tyr Gln Phe Ile His Ala Ala
 145 150 155 160
 Asp Met Leu Tyr Cys Ala Glu Ser His Ile Arg Met Ile Lys Thr Gly
 165 170 175
 Glu Ser Gly Met Ile Val Phe Arg Leu Leu Thr Lys Asn Asn Arg Trp
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 Thr Trp Val Gln Ser Asn Ala Arg Leu Leu Tyr Lys Asn Gly Arg Pro
 195 200 205

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 1 5 10 15
 Gly Val Asp Glu Ala His Gly Pro Pro Gln Ala Ala Val Tyr Tyr Thr
 20 25 30
 Pro Asp Gln Leu Pro Pro Glu Asn Ala Ser Phe Met Glu Arg Cys Phe
 35 40 45
 Arg Cys Arg Leu Arg Cys Leu Leu Asp Asn Ser Ser Gly Phe Leu Ala
 50 55 60
 Met Asn Phe Gln Gly Arg Leu Lys Tyr Leu His Gly Gln Asn Lys Lys
 65 70 75 80
 Gly Lys Asp Gly Ala Leu Leu Pro Pro Gln Leu Ala Leu Phe Ala Ile
 85 90 95
 Ala Thr Pro Leu Gln Pro Pro Ser Ile Leu Glu Ile Arg Thr Lys Asn
 100 105 110
 Phe Ile Phe Arg Thr Lys His Lys Leu Asp Phe Thr Pro Ile Gly Cys
 115 120 125

Asp	Ala	Lys	Gly	Gln	Leu	Ile	Leu	Gly	Tyr	Thr	Glu	Val	Glu	Leu	Cys
130						135					140				
Thr	Arg	Gly	Ser	Gly	Tyr	Gln	Phe	Ile	His	Ala	Ala	Asp	Ile	Leu	His
145					150					155					160
Cys	Ala	Glu	Ser	His	Ile	Arg	Met	Ile	Lys	Thr	Gly	Glu	Ser	Gly	Met
				165					170					175	
Thr	Val	Phe	Arg	Leu	Leu	Ala	Lys	His	Ser	Arg	Trp	Arg	Trp	Val	Gln
			180					185					190		
Ser	Asn	Ala	Arg	Leu	Ile	Tyr	Arg	Asn	Gly	Arg	Pro				
	195						200								